

BRS L1	2 20030082800.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L2	2 6054299.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L3	2 5436141.pn.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L4	120 ssDNA with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L5	3734 (single adj strand adj dna) or ssDNA with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L6	159 ((single adj strand adj dna) or ssDNA) with expression	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L7	47 reverse adj transcriptase and l6	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT
BRS L8	4 reverse adj transcriptase and l6 and retron	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT

L6 ANSWER 1 OF 1 REGISTRY COPYRIGHT 2007 ACS on STN
RN 336761-66-1 REGISTRY
CN DNA (Corynebacterium glutamicum strain ATCC13032 clone
EP1108790-Sequence-3471 gene) (9CI) (CA INDEX NAME)
OTHER NAMES:
CN 3470: PN: EP1108790 SEQID: 3471 claimed DNA
FS NUCLEIC ACID SEQUENCE
SQL 1230
NA 282 a 346 c 352 g 250 t

PATENT ANNOTATIONS (PNTE):

Sequence | Patent
Source | Reference

Not Given | EP1108790

|claimed SEQID
|3471

SEQ	1 atgcttaggca tgcttcgatg gactacagca ggtgaatccc acggccaggc 51 gcttatacgcc acgggttgaac acatgccagc aggctgtgccc gtgactaaag 101 atgaggtctc gtatcaattt ggcgcggccac gccttggata tggctcgccgc 151 gctcgcatga agttttagca agacgcgtt accttcctca cccggcatcccg 201 ccacggcctc actttggta gccccatctc aatcatgtatc ggcaacactg 251 agtgggataa gtggaccacc atcatgtctt ctgacgtttt ggacatggaa 301 gaccaggata acgttgcggc gatgtcttc ggtcgggcgcaaaaactgtac 351 tcgtccgcgtt ccagggccacg ctgattacgc aggcatgttc aagtacgat 401 tcgatgtatgc ccgcaacgtt ctggagcggtt cttagccccgtgaaacactt 451 gcacgcgtgg cagcagcaac ctgtgcgcgt tccttcctgc 501 gggcgtggaa gtgttttttttcc acgttaatttc cattttgtgcg
	===== ===== 551 acactggcgc ggagccaaacc tttgcagata ttcaagcaat cgatgattcc 601 ccagttcgtt catttcgtt aaagacgttggaa gaatccatgtc tgccggaaat 651 cgaggccgcac aagaagccg gcgataccctt cgggtggcatc gtggaaagtga 701 ttgttgaagg cctgcccattt gttttgggtt ctttttttttcc 751 cgcctcgatgtt cgcagatcgcc agctgcactt atgggcatttcc 801 gggcgtggaa atcgggtgacg gtttgcgaa agctcgatcgatcc 851 aagccacgatgtt tgaagtgttcc ctggatgacca acggcgatcc 901 aaccgtgcacg gtggcctcgatcc agggccggatcc accaaccgttcc 951 cgttcgtgttcc ggcatttttttcc caatttttttcc 1001 ccattgtatgttcc ggaaaacggc aaggcagccaa ccggaaatcc 1051 gacgtgtgttcc ctgtttccatcc cggccgggttcc 1101 cctgggttccatcc gcccgcgcacg ttttttttttcc 1151 gcgaaacccaa gagcaacattt gacacccatcc tcaaaaacatcc 1201 atgaaatttcc aagggtttaga ggttttttttcc

HITS AT: 534-546

RELATED SEQUENCES AVAILABLE WITH SEQLINK

MF Unspecified

CI MAN

SR CA

LC STN Files: CA, CAPLUS, TOXCENTER, USPATFULL

DT.CA CAplus document type: Patent

RL.P Roles from patents: BIOL (Biological study); OCCU (Occurrence); PRP (Properties)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH, WPIDS' ENTERED AT 14:15:43 ON
16 APR 2007

L1 9094 S (SINGLE(W) STRAND) (2W) DNA OR SSDNA WITH EXPRESS?
L2 11010 S (SINGLE(W) STRAND) (2W) DNA OR SSDNA(L) EXPRESS?
L3 2957 S ((SINGLE(W) STRAND) (2W) DNA OR SSDNA) (L) EXPRESS?
L4 0 S L3 AND REVERSE (W) TRANSSCRIPTASE
L5 85 S L3 AND REVERSE (W) TRANSCRIPTASE
L6 67 DUPLICATE REMOVE L5 (18 DUPLICATES REMOVED)

(FILE 'HOME' ENTERED AT 16:47:51 ON 16 AUG 2007)

FILE 'REGISTRY' ENTERED AT 16:48:10 ON 16 AUG 2007

L1 3 S TGGTGCCTCCGAG/SQEN
L2 3 SEQLINK EXACT L1

FILE 'CAPLUS' ENTERED AT 16:50:44 ON 16 AUG 2007

L3 3 S L2
L4 3 S L1
L5 0 S L4 NOT L3

FILE 'REGISTRY' ENTERED AT 16:59:11 ON 16 AUG 2007

L6 3 S GTTTCGAAGGCTAGCTACAAACGATCATCCAG/SQEN

FILE 'CAPLUS' ENTERED AT 16:59:59 ON 16 AUG 2007

L7 3 S L6
L8 3 S L7 NOT LL3
L9 1 S L7 NOT L3

FILE 'REGISTRY' ENTERED AT 17:03:55 ON 16 AUG 2007

L10 29 S GGCTAGCTACAAACGA/SQEN

SEQID 7

FILE 'REGISTRY' ENTERED AT 17:04:24 ON 16 AUG 2007

L11 29 S L10 NOT L3 NOT L7